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1600

#15
J. M. M.
12/21/02

RAW SEQUENCE LISTING

DATE: 12/13/2002

PATENT APPLICATION: US/09/674,195B

TIME: 13:54:12

Input Set : A:\W159565.txt

Output Set: N:\CRF4\12132002\I674195B.raw

```
4 <110> APPLICANT: Rosely M. Zancope-Oliveira
5      Timothy J. Lott
6      Leonard W. Mayer
7      Errol Reiss
8      George S. Deepe
11 <120> TITLE OF INVENTION: NUCLEIC ACIDS OF THE M ANTIGEN GENE OF
12      HISTOPLASMA CAPSULATUM, ANTIGENS, VACCINES, AND ANTIBODIES,
13      METHODS AND KITS FOR DETECTING HISTOPLASMOSIS
16 <130> FILE REFERENCE: 14114.0325U2
18 <140> CURRENT APPLICATION NUMBER: 09/674,195B
19 <141> CURRENT FILING DATE: 2000-10-26
21 <150> PRIOR APPLICATION NUMBER: PCT/US99/09151
22 <151> PRIOR FILING DATE: 1999-04-27
24 <150> PRIOR APPLICATION NUMBER: 60/083,676
25 <151> PRIOR FILING DATE: 1998-04-30
27 <160> NUMBER OF SEQ ID NOS: 13
29 <170> SOFTWARE: FastSEQ for Windows Version 4.0
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32 <211> LENGTH: 3862
33 <212> TYPE: DNA
34 <213> ORGANISM: Histoplasma capsulatum
36 <220> FEATURE:
37 <221> NAME/KEY: misc_feature
38 <222> LOCATION: 3258
39 <223> OTHER INFORMATION: n = g, a, c or t(u)
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44 ttttattttc attcatcttc tctgtggcaa acatgcaggt atgcgagctc tggaccctgg      180
45 ggtgtggccc ttgatgcata tggtttattt atagccgccc ggaagccctg gcctgttaaa      240
46 ttttggacct cctcccgcga ttctttccaa acttcgtgcg tccgtttccc atttcccccc      300
47 tccccatttg ggttccctat aggccactgc gtgctccact caagaagggt cccagtcaat      360
48 ttgggtcccta ccctctccaa cactatctgc atatgtaata tatatcgata tctaactgcc      420
49 attgattatt tgtcttcttc agcatctttt tgtctcgagc aagcttactc cacgttcaat      480
50 tcaggggggta aaaatgcggt cgctcaagct tatactcgcc tcggcggggtg ttgtttctgc      540
51 agcctgtccc tacatgtcag gggagatgcc tagcggtcag aaaggccccc tcgatcgccg      600
52 ccattgacact ctctccgacc ctacggacca gtttcttagc aagttttaca ttgacgatga      660
53 acagtcggtg ctaacaacgg acgtgggtgg tcccatcgag gaccaacaca gcctgaaggc      720
54 tggaaaataga ggcccaactc tacttgagga ttttatcttc cgccagaaga ttcaacactt      780
55 tgatcatgag agggtatgta gatacaaaat atgtgaccgt gttgcaaadc cgctaattca      840
56 attttacgca gggtccctgag cgcgcggtcc atgctcgagg agctgggtgcc catggcgat      900
57 tcacatccta taataactgg tcgaatatca cagccgcata cttcttgaac gcggcaggaa      960
58 agcagacacc agtattcgtg cggttttcta cagtcgctgg tagcagaggc agtggttgact      1020
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61 acctaggctg actactcggc aatgtagata tcgtcggaaa caacggtcca gtcttcttca 1200
62 ttcaggacgc tattcaattc cctgatttga ttcacgctgt caagccgcaa ccagacagt 1260
63 aaattcccca ggctgcaact gcacatgata cggcatggga tttcctcagc cagcagccca 1320
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65 atgttgatgg gtggggcgct cataccttcc gacttgtcac cgacgagggc aactcgacct 1440
66 tggtaagtt tcgctggaag accctccaag gaagagcggg cctggtatgg gaagaggcac 1500
67 aggctcttgg cggaaagaat cccgacttcc atcgacaaga cctctgggat gccattgaat 1560
68 ctggaaggta ccctgagtgg gaggtaagat atgattcccc caaatcatta gttctgacag 1620
69 tgtttctctg ctctgtcggg tgctcttttc gtctttttct atatcttcaa ctaagactga 1680
70 ctttatatac gttttactca tatagctggg ctttcaattg gtgaatgaag cagatcaatc 1740
71 caagtttgat ttcgatctat tagatccac aaaaatcatc ccagaagaac ttgttccttt 1800
72 caccccaatc ggaaaaatgg tcttgaaccg aaacccaaaa agttattttg ccgaaactga 1860
73 gcagatcatg gttggtccac cccctatata tttggaatat gaatacatgt atagctagat 1920
74 gaagcgtata tctaaatata tttccacagt tccaaccagg tcatgtagtt cgcggaatcg 1980
75 atttcacgga tgacctttg cttcagggcc gcttgactc ctaccttgac actcaattga 2040
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85 ctaccacaac aaggcaaccg tccccatcgg caccttcggc acgaatctcc tgcggctcga 2640
86 cgggtgaaa atcgccctcc tgacaagaga cgacgtagc ttcacgatcg cggagcagct 2700
87 ccgggcccgc tttaacagcg ccaacaacaa agtagatatc gtcctagtgg gctcatcgct 2760
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95 ctgtttttcc atctttggtt gaggtaatat tgcagatatc agtaaattgc gtttacgaaa 3240
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98 tctcacattc aatctatatt tgatccttgt cctttgtagc tgtttaaata taagccaaga 3420
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104 atagataaaa cctgaacgat ttgacccctg ttgggggaaa taggggttag gggggcgagc 3780
105 tacatatcat tcccatatga ccaaaaaacta aaatagatat atatatatat atatatatat 3840
106 acaacacctt caaaaaggat cc 3862
109 <210> SEQ ID NO: 2

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110 <211> LENGTH: 707
111 <212> TYPE: PRT
112 <213> ORGANISM: Histoplasma capsulatum
114 <400> SEQUENCE: 2
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116 1 5 10 15
117 Ser Asp Pro Thr Asp Gln Phe Leu Ser Lys Phe Tyr Ile Asp Asp Glu
118 20 25 30
119 Gln Ser Val Leu Thr Thr Asp Val Gly Gly Pro Ile Glu Asp Gln His
120 35 40 45
121 Ser Leu Lys Ala Gly Asn Arg Gly Pro Thr Leu Leu Glu Asp Phe Ile
122 50 55 60
123 Phe Arg Gln Lys Ile Gln His Phe Asp His Glu Arg Val Pro Glu Arg
124 65 70 75 80
125 Ala Val His Ala Arg Gly Ala Gly Ala His Gly Val Phe Thr Ser Tyr
126 85 90 95
127 Asn Asn Trp Ser Asn Ile Thr Ala Ala Ser Phe Leu Asn Ala Ala Gly
128 100 105 110
129 Lys Gln Thr Pro Val Phe Val Arg Phe Ser Thr Val Ala Gly Ser Arg
130 115 120 125
131 Gly Ser Val Asp Ser Ala Arg Asp Ile His Gly Phe Ala Thr Arg Leu
132 130 135 140
133 Tyr Thr Asp Glu Gly Asn Phe Asp Ile Val Gly Asn Asn Val Pro Val
134 145 150 155 160
135 Phe Phe Ile Gln Asp Ala Ile Gln Phe Pro Asp Leu Ile His Ala Val
136 165 170 175
137 Lys Pro Gln Pro Asp Ser Glu Ile Pro Gln Ala Ala Thr Ala His Asp
138 180 185 190
139 Thr Ala Trp Asp Phe Leu Ser Gln Gln Pro Ser Ser Leu His Ala Leu
140 195 200 205
141 Phe Trp Ala Met Ser Gly His Gly Ile Pro Arg Ser Met Arg His Val
142 210 215 220
143 Asp Gly Trp Gly Val His Thr Phe Arg Leu Val Thr Asp Glu Gly Asn
144 225 230 235 240
145 Ser Thr Leu Val Lys Phe Arg Trp Lys Thr Leu Gln Gly Arg Ala Gly
146 245 250 255
147 Leu Val Trp Glu Glu Ala Gln Ala Leu Gly Gly Lys Asn Pro Asp Phe
148 260 265 270
149 His Arg Gln Asp Leu Trp Asp Ala Ile Glu Ser Gly Arg Tyr Pro Glu
150 275 280 285
151 Trp Glu Leu Gly Phe Gln Leu Val Asn Glu Ala Asp Gln Ser Lys Phe
152 290 295 300
153 Asp Phe Asp Leu Leu Asp Pro Thr Lys Ile Ile Pro Glu Glu Leu Val
154 305 310 315 320
155 Pro Phe Thr Pro Ile Gly Lys Met Val Leu Asn Arg Asn Pro Lys Ser
156 325 330 335
157 Tyr Phe Ala Glu Thr Glu Gln Ile Met Phe Gln Pro Gly His Val Val
158 340 345 350
159 Arg Gly Ile Asp Phe Thr Asp Asp Pro Leu Leu Gln Gly Arg Leu Tyr

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160          355          360          365
162 Ser Tyr Leu Asp Thr Gln Leu Asn Arg His Gly Gly Pro Asn Phe Glu
163          370          375          380
164 Gln Leu Pro Ile Asn Arg Pro Arg Ile Pro Phe His Asn Asn Asn Arg
165 385          390          395          400
166 Asp Gly Ala Gly Gln Met Phe Ile Pro Leu Asn Thr Ala Ala Tyr Thr
167          405          410          415
168 Pro Asn Ser Met Ser Asn Gly Phe Pro Gln Gln Ala Asn Arg Thr His
169          420          425          430
170 Asn Arg Gly Phe Phe Thr Ala Pro Gly Arg Met Val Asn Gly Pro Leu
171          435          440          445
172 Val Arg Glu Leu Ser Pro Ser Phe Asn Asp Val Trp Ser Gln Pro Arg
173          450          455          460
174 Leu Phe Tyr Asn Ser Leu Thr Val Phe Glu Lys Gln Phe Leu Val Asn
175 465          470          475          480
176 Ala Met Arg Phe Glu Asn Ser His Val Arg Ser Glu Thr Val Arg Lys
177          485          490          495
178 Asn Val Ile Ile Gln Leu Asn Arg Val Asp Asn Asp Leu Ala Arg Arg
179          500          505          510
180 Val Ala Leu Ala Ile Gly Val Glu Pro Pro Ser Pro Asp Pro Thr Phe
181          515          520          525
182 Tyr His Asn Lys Ala Thr Val Pro Ile Gly Thr Phe Gly Thr Asn Leu
183          530          535          540
184 Leu Arg Leu Asp Gly Leu Lys Ile Ala Leu Leu Thr Arg Asp Asp Gly
185 545          550          555          560
186 Ser Phe Thr Ile Ala Glu Gln Leu Arg Ala Ala Phe Asn Ser Ala Asn
187          565          570          575
188 Asn Lys Val Asp Ile Val Leu Val Gly Ser Ser Leu Asp Pro Gln Arg
189          580          585          590
190 Gly Val Asn Met Thr Tyr Ser Gly Ala Asp Gly Ser Ile Phe Asp Ala
191          595          600          605
192 Val Ile Val Val Gly Gly Leu Leu Thr Ser Ala Ser Thr Gln Tyr Pro
193          610          615          620
194 Arg Gly Arg Pro Leu Arg Ile Ile Thr Asp Ala Tyr Ala Tyr Gly Lys
195 625          630          635          640
196 Pro Val Gly Ala Val Gly Asp Gly Ser Asn Glu Ala Leu Arg Asp Val
197          645          650          655
198 Leu Met Ala Ala Gly Gly Asp Ala Ser Asn Gly Leu Asp Gln Pro Gly
199          660          665          670
200 Val Tyr Ile Ser Asn Asp Val Ser Glu Ala Tyr Val Arg Ser Val Leu
201          675          680          685
202 Asp Gly Leu Thr Ala Tyr Arg Phe Leu Asn Arg Phe Pro Leu Asp Arg
203          690          695          700
204 Ser Leu Val
205 705
208 <210> SEQ ID NO: 3
209 <211> LENGTH: 8
210 <212> TYPE: PRT
211 <213> ORGANISM: Histoplasma capsulatum

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213 <400> SEQUENCE: 3
214 Ser Asp Pro Thr Asp Gln Phe Leu
215 1 5
217 <210> SEQ ID NO: 4
218 <211> LENGTH: 15
219 <212> TYPE: PRT
220 <213> ORGANISM: Histoplasma capsulatum
222 <400> SEQUENCE: 4
223 Asp Phe Ile Phe Arg Gln Lys Ile Gln His Phe Asp His Glu Arg
224 1 5 10 15
226 <210> SEQ ID NO: 5
227 <211> LENGTH: 9
228 <212> TYPE: PRT
229 <213> ORGANISM: Histoplasma capsulatum
231 <400> SEQUENCE: 5
232 Thr Leu Gln Gly Arg Ala Gly Leu Val
233 1 5
235 <210> SEQ ID NO: 6
236 <211> LENGTH: 16
237 <212> TYPE: PRT
238 <213> ORGANISM: Histoplasma capsulatum
240 <400> SEQUENCE: 6
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242 1 5 10 15
244 <210> SEQ ID NO: 7
245 <211> LENGTH: 6
246 <212> TYPE: PRT
247 <213> ORGANISM: Histoplasma capsulatum
249 <400> SEQUENCE: 7
250 Ser Gly Arg Tyr Pro Glu
251 1 5
253 <210> SEQ ID NO: 8
254 <211> LENGTH: 10
255 <212> TYPE: PRT
256 <213> ORGANISM: Histoplasma capsulatum
258 <400> SEQUENCE: 8
259 Phe Asp Phe Asp Leu Leu Asp Pro Thr Lys
260 1 5 10
262 <210> SEQ ID NO: 9
263 <211> LENGTH: 14
264 <212> TYPE: PRT
265 <213> ORGANISM: Artificial Sequence
267 <220> FEATURE:
268 <223> OTHER INFORMATION: Description of Artificial Sequence; M antigen-specific
oligonucleotide
270 <400> SEQUENCE: 9
271 Ile Ile Pro Glu Glu Leu Val Pro Phe Thr Pro Ile Gly Lys
272 1 5 10
275 <210> SEQ ID NO: 10
276 <211> LENGTH: 15

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RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/09/674,195B

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. 3258

Seq#:11; N Pos. 3,9

VERIFICATION SUMMARY

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L:96 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:3240

L:326 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:0